

REVIEW



Rho GTPase regulation of reactive oxygen species generation and signalling in platelet function and disease

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ABSTRACT

Platelets are master regulators and effectors of haemostasis with increasingly recognized functions as mediators of inflammation and immune responses. The Rho family of GTPase members Rac1, Cdc42 and RhoA are known to be major components of the intracellular signalling network critical to platelet shape change and morphological dynamics, thus playing a major role in platelet spreading, secretion and thrombus formation. Initially linked to the regulation of actomyosin contraction and lamellipodia formation, recent reports have uncovered non-canonical functions of platelet RhoGTPases in the regulation of reactive oxygen species (ROS), where intrinsically generated ROS modulate platelet function and contribute to thrombus formation. Platelet RhoGTPases orchestrate oxidative processes and cytoskeletal rearrangement in an interconnected manner to regulate intracellular signalling networks underlying platelet activity and thrombus formation. Herein we review our current knowledge of the regulation of platelet ROS generation by RhoGTPases and their relationship with platelet cytoskeletal reorganization, activation and function.

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Introduction

Platelets are anucleate fragments of megakaryocytes that surveil the vasculature for damage to the endothelium as primary cellular mediators of haemostasis [1–3]. Upon detection of endothelial injury or dysfunction, platelets undergo drastic morphological changes and aggregate with other platelets to protect vascular integrity [4,5]. Following platelet adhesion to the endothelium, cytoskeletal reorganization, namely the formation of actin-rich filopodia and lamellipodia, occurs in conjunction with fibrinogen-dependent platelet aggregation and haemostatic plug formation. Here, small GTP-binding proteins of the *Aplysia* Ras Homologous (ARH), or Rho GTPase family play critical roles in platelet responses that ultimately orchestrate key features of platelet activation and thrombus formation [6,7].

Rho GTPases and their regulators orchestrate platelet cytoskeletal dynamics

The Rho GTPases family consists of small GTP-binding proteins that range from 20 to 40 kDa in size and

includes over 20 members divided into classic and atypical members [6]. Classic Rho GTPases such as RhoA, Rac1, and Cdc42 are regulated by Rho-specific guanine nucleotide exchange factors (GEFs), GTPasesactivating proteins (GAPs), and guanine nucleotide dissociation inhibitors (GDIs). The Rho GTPases cycle between active GTP-bound state and inactive GDPbound state, regulated by GEFs that activate GTPases by promoting exchange of GDP for GTP; GAPs that inactivate GTPases by promoting GTP hydrolysis; and GDIs that shuttle inactive GTPases throughout the cell [8]. Interactions between active GTP-bound GTPases and their effector molecules lead to activation of downstream signalling pathways that are crucial for regulating cellular functions including migration, secretion, and spreading. In platelets, the Rho GTPase family members Cdc42, Rac1 and RhoA are known to play critical roles in platelet haemostatic responses [6,7,9].

Studies over the past two decades have specified how Rac1 promotes platelet lamellipodia formation, whereas RhoA stimulates actomyosin contraction underlying platelet shape change as well as clot retraction. In

general, Rho GTPases are not directly regulated by reversible phosphorylation but by a profound phosphorylation of GEFs, GAPs and other Rho regulators that specify the activities of Rho GTPases in contexts of platelet activation[10]. However, a few direct phosphorylation sites have been identified on RhoA (Ser₁₈₈) [11-14], Rac1 (Ser⁷¹) and Cdc42 (Ser¹⁸⁵, Ser⁷¹, Tyr⁶⁴) [15-17], which have been reported to exert inhibitory effects on Rho GTPase function in some cell types. For instance, direct phosphorylation was first described for RhoA at Ser¹⁸⁸ by protein kinase A (PKA) as part of the cyclic AMP (cAMP) signalling pathway, resulting in increased RhoA-RhoGDI interaction that inhibits RhoA membrane relocalization and therefore reducing RhoA activity [11-14,18]. Furthermore, Cdc42 is also phosphorylated by PKA at Ser¹⁸⁵ and by Src kinase at Tyr⁶⁴, resulting in enhanced interaction with RhoGDI [19,20]. RhoA downregulation by cAMP-PKA signalling pathway has indeed recently been confirmed in platelets, demonstrating that RhoA Ser¹⁸⁸ phosphorylation by PKA prevents the association of RhoA with Rho-associated coiled-coil containing protein kinase (ROCK)2 and myosin phosphatase targeting subunit 1 (MYPT1) in regulating platelet function[15]. On the other hand, Rac1/Cdc42 phosphorylation at Ser⁷¹ by AKT, resulting in reduced GTP-binding without affecting GTPase activity [16,17,21], has yet to be demonstrated in platelets. Overall, it seems that direct phosphorylation does not necessarily inactivate/activate Rho GTPases, but rather modulates their subcellular locations and affinity to Rho GDIs in platelets. Meanwhile, phosphorylation of GEFs, GAPs and Rho regulators governs Rho GTPases activity, especially in platelet activation programs.

Other covalent modifications of Rho GTPases, including oxidation, can directly shape Rho GTPase activities in a manner relevant to platelet function in health and disease. In this review, we highlight the studies defining the mechanisms underlying platelet ROS production and secretion and put into perspective the contribution of Rho GTPases in mediating platelet oxidative stress (Figure 1).

Rho GTPases regulate cellular oxidative stress via redox-sensitive motifs

Intracellular reactive oxygen species (ROS) include free radicals, such as superoxide radical anion and hydroxyl anion, hydrogen peroxide, singlet oxygen, peroxynitrite and hypochlorous acid[22]. Table 1 summarizes platelet-derived ROS and the mechanisms responsible for ROS production. Not all ROS act as cellular messengers due to their short half-life in tissues as a result of spontaneous dismutation or dismutation catalysed by superoxide dismutase[23]. The half-life of hydrogen peroxide is a few seconds, and together with the ability to dissolve in lipids and pass through membranes makes hydrogen peroxide the most likely candidate for intracellular and intercellular signalling[24]. On the other hand, superoxide radical and peroxynitrite function solely as intracellular messengers, acting on proteins in the mitochondrial membrane, including nicotinamide adenine dinucleotide phosphate (NADPH) oxidase, xanthine oxidase and nitric oxide synthase. ROS also act through post-translational modification of receptors, kinases, phosphatases, ion channels and transcription factors, among which the most well known is the hydrogen peroxide-mediated oxidation of cysteine residues in proteins[25].

In addition to regulating cytoskeletal dynamics and vesicular traffic in platelets and other cells, many of the Ras superfamily of small GTPases with GXXXXGK(S/ T) motifs, including Rho GTPases, also participate in redox-sensitive regulatory processes [26,27]. In a manner similar to GEFs, redox-sensitive motifs allow perturbation in nucleotide-binding interactions, resulting in enhanced guanine nucleotide exchange of GTPases [26]. Historically, ROS and reactive nitrogen species are considered mainly as by-products of aerobic metabolism and other enzymatic processes whose upregulation in turn imposes toxicity to macromolecules including DNA, proteins and lipids. However, ROS formation is now recognized as a tightly regulated process involved in multiple cellular signalling pathways related to cell adhesion[28], growth factor signalling[29], inflammation and host defence[30]. ROS-mediated oxidation of thiol groups can modulate the activity of proteins that rely on cysteine fragments for substrate binding such as p38 kinase, tropomyosin and many of the tyrosine [31,32]. Ca²⁺/calmodulin-dependent phosphatases kinase II[33], protein kinase G (PKG)[34], protein kinase A (PKA)[35], extracellular signal-regulated kinase (ERK)[36], phosphoinositide 3-kinase (PI3K), Akt[37], protein kinase C (PKC)[38],N-terminal kinase [39] are among many targets of ROS that are critical for intracellular signalling. Rho GTPases, ROS and cytoskeletal organization function as an interconnected regulatory network that drives intracellular signalling underlying thrombus formation and cellular oxidative stress, although the precise means of regulation of ROS production by Rho GTPases is not completely understood [26,40–43].

While initially it was thought that endothelial cells, leukocytes, smooth muscle cells and fibroblasts act collectively as the main source of ROS, recent studies have shown that activated platelets act as an additional and

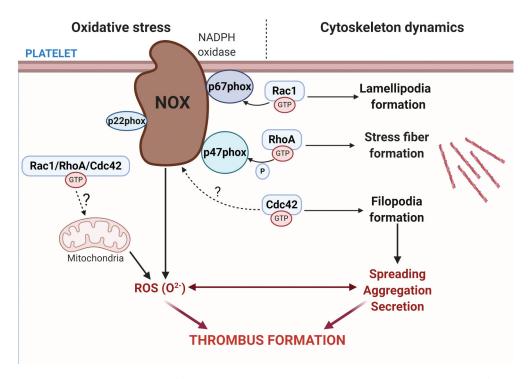


Figure 1. Platelet RhoGTPases regulate ROS formation. Platelet express the most well-known RhoGTPases family members Rac1, Cdc42 and RhoA that regulates actin reorganization within the platelet cytoskeleton. Rac1 and Cdc42 regulate lamellipodia and filopodia formation, while RhoA regulates stress fiber formation. In addition to these classical roles in platelet activity, RhoGTPases have been uncovered as regulators of reactive oxygen species (ROS) formation by modulating the NADPH oxidase complex. Rac1 regulates NADPH oxidase complex assembly by directly binding to p67^{phox} in response to thrombin, as well as GPVI and GPIb agonists. RhoA regulates thrombin-mediated ROS formation by inhibiting p47^{phox} phosphorylation, whereas a potential role for Cdc42 in directly regulating ROS formation remains to be identified. ROS generation triggers a positive feedback loop on platelet activation that promotes further ROS production and amplifies platelet recruitment, activation and aggregation, which ultimately leads to thrombus formation.

potentially leading source of superoxide anion, hydroxyl radicals and hydrogen peroxide, due to the biomass of platelets relative to other blood cells [44-48]. Vascular cells contain numerous potential sources for ROS generation including mitochondrial electron transport chain, lipoxygenase, xanthine oxidase, cyclooxygenase and the NADH/NADPH oxidase system[49]. The main source of platelet ROS production is derived from NADPH produced in the pentose cycle [50], while ROS production as a respiratory by-product of mitochondria plays a secondary role[51].

Potential prothrombotic role of ROS

The role of ROS has been implicated in a number of conditions with increased risk for deep vein thrombosis (DVT) such as antiphospholipid syndrome (APS), autoimmune disorder (Behcet's disease) and metabolic syndrome among others [52]. For instance, in an APS mouse model, cofactor-independent antiphospholipid antibodies induce a procoagulant state leading to venous thrombosis via activation of endosomal NADPH oxidase

[53,54]. Hydroxychloroquine, used to treat patients with APS, targets endosomal NOX[55]. In Behcet's disease, thrombus formation is promoted by oxidized fibrinogen from enhanced NOX-dependent ROS production[56]. Plasma of obese individuals has been shown to contain elevated levels of CD40 ligand and soluble P-selectin, both associated with NOX2 upregulation in platelets [57]. Oxidative stress induces oxidation of low-density lipoproteins, which bind to CD36 and induce platelet activation via NOX2 and may play a role in prothrombotic mechanisms in patients with metabolic syndrome [58–60]. ROS stimulates coagulation by increasing tissue factor expression in endothelial cells, monocytes and smooth muscle cells via NOX enzymes [61-65]. Furthermore, many calcium-based signalling systems in cells interact with redox signalling pathways. NOX enzymes are regulated by calcium, and calcium channels are regulated by oxidative modifications [66-71]. As a result, calcium-ROS interplay underlies processes by which leukocytes can contribute to thrombus formation via NOX2[65]. Platelet activation relies on calcium surges driven by store-operated calcium entry via

Table 1. Platelet-derived reactive oxygen species (ROS) and their mechanisms of production in platelets

Platelet-derived ROS	Enzymes involved and mechanisms of ROS production
Superoxide (O ₂ ⁻)	Upon GPVI activation
	 Via lipoxygenase and the glutathione cycle[117,225] Mediated by NADPH oxidase activity via PI3K pathways[116,226] Arachidonic acid (AA)-dependent[227] Upon stimulation by thrombin
	 Mediated by PKC and PI3K[^{228]} By other agonists
	 Calcium ionophore and tissue plasminogen activator induce O₂⁻ generation in an NAD(P)H-dependent manner[^{50]} Anoxia-reoxygenation triggers intrinsic production of O₂^{-[109]} Bacterial LPS induces O₂⁻ generation mediated by PKC and Pl3K[^{228]}
Hydrogen peroxide (H ₂ O ₂₎	Upon GPVI activation
	 Collagen (but not thrombin or ADP) induces H₂O₂ production[^{125]} By other ligands
	 Zymosan and latex beads trigger H₂O₂ production in an NADH-dependent manner[^{229]}
Hydroxyl (OH ⁻)	Upon GPVI activation
	 Via AA-dependent mechanism[^{227]} By other agonists
	 Anoxia-reoxygenation triggers intrinsic production of OH⁻¹⁰⁹ AA triggers OH⁻ formation, likely through the lipoxygenase pathway[^{230]} Iron directly activates platelets via OH⁻ formation, mediated by PKC activity[^{231]}
Unspecified ROS	Upon GPVI activation
	 Mediated by ITAM signalling via Syk-dependent and independent pathways[^{128]} TRAF4 links GPVI with p47^{phox} as a mean to promote ROS formation[^{127]} <i>Upon stimulation by thrombin</i>
	 Downstream of GPIb_α and PAR4 → ROS formation through FAK and NADPH oxidase-dependent mechanisms[^{144,232]} Via xantine oxidase, PKC and PI3K-dependent pathways[^{44]} By other agonists
	 Zn²⁺ regulates ROS generation in an NADPH oxidase and mitochondria activation-dependent manner, mediated by MAPK pathways[^{233]} VPS34 is a major mediator of platelet activation via NADPH oxidase-dependent ROS generation[^{234]} Hyperglycaemia increases ROS generation in human platelets via mitochondrial electron transport system-derived ROS[^{76]} Hyperthermia promotes platelet mitochondrial ROS formation in a time-dependent manner[^{235]} LDL oxidation by platelets subsequently amplifies platelet activation mediated by NOX2 activation [^{236]}

stromal interaction molecule 1 protein and plasma membrane channel Orai1, which are under redox control [66,67,72].

In platelets, the generation of ROS regulates signalling pathways involved in platelet activation and thrombus formation, as overproduction of ROS is associated with hypercholesterolemia, diabetes, hypertension, and metabolic syndrome, all of which are risk factors for thrombosis. Therefore, ROS generation may play a prothrombotic role in select diseases, especially those with associated risk for thrombosis. However, a casual link between platelet Rho GTPase-mediated ROS signalling and thrombosis *in vivo* remains to be specified.

Mitochondria-mediated ROS formation in platelets

Platelets do not contain nuclei [73] and their lifespan of 5–7 days in circulation depends heavily on the mitochondria, which are critical for aerobic respiration, and also yield metabolic substrates needed for platelet function at rest and during energetically-demanding processes such as activation, secretion and aggregation [74–76]. Discordant studies debate whether glycolysis or oxidative phosphorylation constitute the major source of ATP in platelets [77,78], as inhibition of either metabolic process abrogates platelet aggregation [79]. Indeed, a recent study has demonstrated that platelets exhibit metabolic plasticity depending on the

substrates used for aggregation[80]. Mitochondria link the energy-releasing activities of electron transport with the process of oxidative phosphorylation to harness fuel in the form of ATP. Mitochondria are both effectors and primary targets of cardioprotective signalling in a range of cells, including platelets[38]. In patients and animal models of sepsis, the mitochondria are swollen and their matrix is disrupted, associating with a decrease in cellular oxidative capacity[81]. Opening of the mitochondrial permeability transition is influenced by high levels of calcium, phosphate and ROS, and is responsible for the pathogenesis of necrosis following ischaemia-reperfusion and organ dysfunction accompanying septic shock [82-84].

Mitochondrial damage or dysfunction results in attenuated platelet survival and increased risk for thrombosis. 'Leaky' reactions of the electron transport chain with molecular oxygen lead to the generation of superoxide anion radicals that subsequently forms hydrogen peroxide and hydroxyl radicals[85]. As one of the drivers of this process, the serine protease and coagulation factor thrombin induce mitochondrial depolarization, cytochrome c release, activation of caspases-3 and -9, and phosphatidylserine exposure in platelets[86]. This is particularly relevant as the platelet surface and secretome acts as a catalyst for thrombin generation, thus creating a potential feedback mechanism by which platelets can incite mitochondrial-mediated ROS formation. Mitochondrial permeability transition pore (mPTP) openings, together with inner membrane anion channel (IMAC) play an important role in the mitochondrial adaptive responses to redox stress [87,88]. Mitochondrial ROS production involves a biphasic mechanism (ROS-induced ROS release), wherein oxidative challenge results in an amplified ROS signal, which depending on ROS levels may result in different outcomes [89-93]. Reversible mPTP opening and/or IMAC-associated ROS release acts as a housekeeping function by allowing the timely release of accumulated toxic levels of ROS. As inner ROS levels continue to rise, longer mPTP openings may release a ROS burst, ultimately leading to mitochondrial destruction as a physiological response to remove unwanted cells and damaged mitochondria, or as a pathological response leading to the elimination of vital and essential mitochondria [48,89-93]. ROS released into the cytosol could trigger signalling responses and ROS-induced ROS release in neighbouring mitochondria. As a consequence, ROS, generated by the mitochondria and at other sites, induces cytotoxicity resulting in platelet oxidative stress, thrombocytopenia and bleeding[94].

In rabbit synovial fibroblasts, clustering of integrins by anti-α5 integrin antibodies (integrin ligation) leads to changes in cell adhesion followed by actin cytoskeletal reorganization as well as ROS production and subsequent nuclear factor (NF)-κB activation via a Rac-dependent mechanism[95]. In this context, Rac activation occurs downstream of integrin crosslinking and prior to ROS production, where inhibition of mitochondrial respiratory chain complexes by rotenone, antimycin or potassium cyanide abrogates hydrogen peroxide generation. These and other observations indicate that mitochondria are the main source of integrin-mediated ROS production in fibroblasts via Rac activity[96]. A few mechanisms have been suggested for Rho GTPases in modulating mitochondrial oxidative function. One involves the control of the anti/ proapoptotic function of the B-cell lymphoma (Bcl)-2 family members, together with the organization of the cell cytoskeleton. Rac also activates PI3K and Akt, which modulates the apoptotic function of Bcl-2-associated agonist of cell death (Bad). However, inhibition of PI3K did not affect integrin ligation-induced ROS potential mechanisms formation[96]. Additional include an interaction of Bcl-2 with Rac and p190RhoGEF during cell spreading. Corroborating this, Velaithan et al. later on showed direct physical interaction between mitochondrial Rac1 and Bcl-2 in human cancer cell lines, maintaining a pro-oxidant intracellular milieu through increased ROS levels[97]. Furthermore, increased mitochondrial Rac1 activity in alveolar macrophages via cytochrome c increased oxidative stress associated with pulmonary fibrosis[98], and was proposed to play important roles in neuroplasticity and anti-apoptosis and autophagy via inositol 1,4,5-trisphosphate receptor and Bcl-2 at the mitochondrial membrane[99].

In other cell types, including platelets, p38 phosphorylates and activates the mitogen-activated protein kinase (MAPK) associated kinase MK2 in response to cellular stresses leading to phosphorylation of MK2 substrates such as RTN4 and Bcl-2 family member Bcl-xl[100]. Although most studies have focused on how Bcl-2 proteins regulate mitochondrial outer membrane during apoptosis [101,102], growing evidence also indicates that Bcl-2 proteins are localized and function on the endoplasmic reticulum (ER), with distinct roles for mitochondrial and ER-localized Bcl-xl [103–105]. While mitochondrial Bcl-xl regulates apoptosis independent of ER Bcl-xl, the involvement of ER Bcl-xl in antiapoptotic signalling is rather linked to its role in calcium homoeostasis [106,107]. Therefore, in response to thrombin stimulation, platelets upregulate Bcl-xl phosphorylation as an effort to organize the ER relative to the mitochondria to facilitate calcium signalling events necessary for platelet activation. Given the roles of p38-Bcl-xl axis in mediating platelet phosphatidylserine exposure through apoptosis-related pathways, future efforts placing Rho GTPase-mediated oxidative stress and ROS production into the context of platelet activation programs may help to better understand the role of Rho GTPase signalling rooted in the ER-mitochondria communication.

NADPH oxidase-mediated ROS formation in platelets

The NADH/NADPH oxidase system was originally described as a primary source of ROS in phagocytic cells. Leoncini et al. were among the first to show that the enzymatic activity of NAD(P)H-cytochrome c reductase was present in human platelets[108]. Inhibition of NADPH oxidase suppresses ROS release by platelets and subsequent platelet aggregation under experimental conditions in a mouse model of ischaemia-reperfusion [109,110]. The NAD(P)H components, including phagocyte oxidase (phox) subunits gp91^{phox} (NOX2), p22^{phox}, p40^{phox}, p47^{phox}, and p67^{phox}, become activated in response to pro-inflammatory mediators[111]. The best characterized member of the NOX family is mammalian NOX2 (gp91^{phox}), which is the most highly expressed member in phagocytes such as neutrophils[112]. Six homologs of NOX2, namely NOX1, NOX3, NOX4, NOX5, DUOX1 and DUOX2 have been characterized in various cell types[113]. Human platelets and megakaryocytes (MEG01 cells) express NOX1[114], NOX2 [114,115], together with cytosolic cofactors p22^{phox}, p47^{phox} and p67^{phox} [50,116,117]. Murine megakaryocytes also express NOX1 and NOX4[118]. The presence of NOX4 [45] and NOX5 [119] in platelets still remains controversial.

Although enzymatic activity of NADPH oxidase is greater in phagocytes as compared to platelets on a per cell basis[50], it is increasingly clear that platelets contribute to diverse immunological processes beyond haemostasis and thrombosis. Accumulating evidence suggests the ability for platelets to directly interact with and phagocytose viral pathogens [120,121], wherein the released ROS from platelets have antimicrobial role and contribute to the killing of pathogens [122]. NOX1 and 2 have been implicated in promoting platelet activation, secretion and aggregation through ROS formation downstream of G-protein-coupled receptors (GPCRs) and glycoprotein [86,114,123,124]. Platelet GPVI-mediated signalling through the NADPH oxidase complex is now thought to be a key mediator of intracellular ROS formation in platelets [117,125,126] via two distinct phases [127]. The initial phase is Syk-independent, with TRAF4 mediating the interaction of Lyn with PKC $_{\delta}$, which in turn phosphorylates p47^{phox} and leads to a rapid burst of ROS. The second phase is Syk-dependent; with ITAM signalling leading to Syk activation and subsequent PLC_v2-IP3-PKC axis activating NOX and supporting further ROS production [47,48,128].

Rac1: 'NOX' your typical first responder

The small GTPase Rac is required for NADPH oxidase activity in both phagocytic and non-phagocytic cells [129,130], with homologs of NOX2 (gp91^{phox}) identified in non-phagocytic cells as well, including platelets [113]. Only Rac, but not Cdc42 or RhoA, is capable of functioning as a direct activator of NOX1-3, where Rac is the most studied Rho GTPase in oxidative stress pathways [128,131]. Upon stimulation, two cytosolic components of NADPH oxidase, p47^{phox} and p67^{phox}, translocate to the plasma membrane and form a complex with NOX2 and p22^{phox}, which are two subunits of a low-potential, membrane-bound flavocytochrome b [129]. Active Rac GTPase translocates to the plasma membrane simultaneously to but independently of the translocation of p47^{phox} and p67^{phox}[132]. Once formed, the NADPH oxidase complex acts as an electron transporter, carrying electrons from NADPH via FAD and then to the haem groups of cytochrome b, directly reducing O₂[133].

Rac1 regulates both NOX2 (phagocyte oxidase) and NOX1 (non-phagocytic oxidase)[134]. NOX2 activation is completely dependent on Rac. The binding of Rac1-GTP to p67^{phox} facilitates its binding to NOX2 and its activation, generating superoxide [135,136]. Similarly, binding of Rac1-GTP to NOXA1, a p67^{phox}related protein, enhances its binding to NOX1 and its activation [41,137]. Interestingly, it has been demonstrated that Rho GDI stabilizes Rac in an active conformation as well, even in the GDP-bound state, and presents Rac to p67^{phox} of the NADPH oxidase complex [42,138-140]. This concept challenged traditional beliefs regarding the role of Rho GDIs in regulating Rho GTPases activity, where Rho GDIs are classically thought of as passive shuttles of Rho GTPases [138,141]. Therefore, Rho GDIs might employ more direct roles in regulating Rho GTPase activity than currently appreciated [8,142].

In platelets, Rac GTPase signalling plays an important role in thrombin-mediated ROS generation and platelet activation. As an example, studies showed that platelets from Rac1^{-/-} conditional knockout mice or human platelets treated with NSC23766, a Rac inhibitor, exhibited diminished thrombin-induced ROS generation[41]. Blocking Rac1-p67^{phox} interaction was demonstrated to inhibit NOX2 activation and ROS formation in human and murine neutrophils as well [143]. While the exact mechanisms of ROS generation in platelets are still being delineated, Carrim et al. demonstrated that a functional cooperation between GPIbα and PAR4 is required for thrombin-induced ROS formation, mediated by NOX1 and focal adhesion kinase. Interestingly, such cooperative role was not shared by the thrombin receptor, PAR1, which is only expressed in human and not mouse platelets[144].

Given its roles in platelets and other cellular mediators if inflammation, Rac has been put forth as a druggable target, although the translation for use in cardiovascular disease would require balancing selectivity, efficacy and safety. In purified platelet systems, inhibition of Rac1-p67^{phox} interaction using a small molecule inhibitor (Phox-I) has been demonstrated to prevent GPVI- and non-GPVI (thrombin receptor)mediated NOX2 activation and subsequent ROS generation, as well as inhibit in vitro and in vivo platelet activation, spreading and aggregation by blocking PI3K activation and downstream phosphorylation of Akt, as well as ERK and p38 MAPK[124]. Akbar et al. first demonstrated that a reversible inhibitor of Racp67^{phox} interaction and therefore inhibition of NOX2 activation was capable of altering platelet function without compromising the haemostatic response[124]. Perhaps pharmacological targeting of NOX2 via Rac GTPase activity may present a new antithrombotic approach by preventing pathological GPVI- and thrombin-mediated NOX2 activation and subsequent ROS generation while preserving physiological platelet functions. Still, selectivity would need to be addressed to ensure the safety of such an approach, seeing that Rac activity plays a plethora of physiological roles in haematopoietic and vascular cells alone. Furthermore, although it is well known that Rac1 is essential for platelet spreading and aggregation [145-147], data still remain conflicting whether NOX2 functionally mediates superoxide generation in platelet activation and thrombosis. For instance, some studies suggest NOX2 oxidase is dispensable for platelet ROS production, while others emphasize the importance of NOX2 (in addition to NOX1) in mediating experimental thrombosis [123,148].

RhoA to the 'ROScue'

The Rho subclass of the Rho GTPases family is composed of three highly conserved proteins: RhoA, RhoB and RhoC. Among them, RhoA has been the most

studied member. Although the three isoforms were discovered contemporarily, proteomic and transcriptomic studies revealed that RhoA is the dominant member expressed in human platelets as compared to RhoB and RhoC [149,150]. Whereas Rac1 and Cdc42 are critically involved in the regulation of lamellipodia and filopodia formation, RhoA regulates actomyosin contractility as well as actin-myosin stress fibre and focal adhesion formation[151]. The signalling events leading to RhoA activation involve the activation of upstream G₁₂ and G₁₃ proteins and downstream Rhoassociated coiled-coil containing protein kinases (ROCK), subsequently promoting myosin light chain (MLC) phosphorylation, which is a major promoter of platelet shape change in both human and mouse platelets [152,153]. Activation of GPCRs by thrombin promotes $G_{\alpha\alpha}$ and p115RhoGEF activation, which triggers the formation of a RhoA-GTP complex and promotes platelet contractile activity as well as granule secretion [154]. In a secondary step of platelet activation, $G_{\alpha\alpha}$ activates c-Src to promote a negative feedback loop that involves p190RhoGAP, stimulating the hydrolysis of RhoA-GTP to RhoA-GDP and therefore diminishing RhoA-induced contraction, which in turn facilitates cell spreading[155].

Changes in the cellular redox state and regulation of the actin cytoskeleton by Rho GTPases are indeed deeply interconnected processes. In conjunction with but independent of the RhoA/ROCK-mediated phosphorylation of MLC that is required for maintenance of platelet structure during spreading and thrombus stability[156], RhoA activation and downstream ROCKmediated activation of p38 MAPK and ERK1/2 also lead to subsequent p47^{phox} phosphorylation, activation of the NADPH oxidase and ROS generation, which further contributes to platelet activation [40,43]. The first evidences for RhoA playing a crucial role in platelet ROS generation were demonstrated in platelets treated with Rhosin, a small molecule inhibitor of RhoA, Y27632, a known inhibitor of RhoA, and in platelets from RhoA^{-/-} mice. RhoA deficiency or pharmacological inhibition of RhoA reduced platelet ROS generation in response to thrombin [40,153,157]. RhoA regulates ROS formation indirectly by inhibiting phosphorylation of p47^{phox}, in contrast to how Rac1 regulates ROS generation through direct Rac1-p67^{phox} interaction [40,124].

Although there has not been any evidence suggesting a direct role of RhoA in NADPH oxidase activation, the redox-mediated regulation of RhoA activity has been implicated in the Rac1-mediated generation of ROS through NADPH oxidase. Several studies have demonstrated that the activity of Rho can be downregulated by Rac and that the actin cytoskeletal reorganization induced by such antagonistic relationship dictates cellular morphology [158,159]. Tiam1/Rac1 signalling antagonizes Rho activity directly. Rac1, but not Rac2 or Rac3, facilitates p190RhoGAP activation by Src kinase-dependent tyrosin phosphorylation or by enhancing p190RhoGAP recruitment to the cell membrane through interaction with p120RasGAP, ultimately driving rapid Tiam1-mediated downregulation of Rho[160]. It is important to note that there are about 60 human RhoGAPs with a common catalytic domain capable of stimulating GTP hydrolysis reaction [161-164]. A recent report has shown that in addition to p190, oligophrenin-1 (OPHN1), a RhoA GAP, is also among RhoGAPs with the highest selectivity and catalytic efficiency towards Rac1[165]. As such, increases in the tyrosine phosphorylation and activation of p190RhoGAP and potentially other RhoGAPs might constitute an integral mechanism for the coupling of changes in cellular redox state to the control of the actin cytoskeleton by Rho GTPases[166].

Studies have also demonstrated alternative mechanisms of RhoA activity regulation independent of classical regulatory proteins such as Rho GAPs, GEFs or GDIs. Some alternative mechanisms entail ROS directly inducing reversible activation of RhoA, mediated by redox-sensitive cysteine residues within the phosphoryl binding loop of RhoA redox-sensitive motif (CXXXCGK(S/T)C), the result of which is stress fibre formation [27,167]. As observed in rat pulmonary artery, ROS, generated via NOX of the NADPH oxidase complex or from mitochondria, activates Src-family kinases via direct oxidation or inhibition of c-Src kinase or inhibitory phosphatases, followed by activation of ARHGEF1, RhoA and ROCK, resulting in enhanced MYPT-1 and MLC₂₀ phosphorylation and cellular contraction[168]. It is yet to be seen whether these mechanisms are conserved and utilized in platelets.

Cdc42: navigator of the actin seas

Rac and Cdc42 signalling, facilitated by the GAP GIT1 and GEFs βPIX and GEFH1, converges on the p21activated kinases (PAK) system and downstream PAK effectors required for thrombin-mediated activation of MEK/ERK pathway, Akt, calcium signalling and phosphatidylserine exposure critical for platelet haemostatic function[169]. Similar to the extended actions of Rho and Rac beyond actin reorganization, Cdc42 activity also functions beyond its classical role in controlling cellular migration through filopodia formation.

The effector domains (Switch I) of Rac1/2 and Cdc42 differ by only four amino acids, however, only

the GTP-bound form of the Rac, but not Cdc42, Switch I domain interacts with p67^{phox}, a direct activator of NADPH oxidase. Mutation of Ala²⁷ and Gly³⁰ residues within the Switch I region enables Cdc42 to function as a direct activator of NADPH oxidase through stabilizing the mutated Cdc42-p67^{phox} complex together with cytochrome b. Although Cdc42 is unable to stimulate ROS formation by directly activating NADPH oxidase, Cdc42 can bind to cytochrome b in vitro and act as a competitive inhibitor of Rac1/2-mediated ROS formation[170]. Neutrophils treated with casin, a Cdc42 activity-specific inhibitor, produce significantly more ROS in response to complement component 5a (C5a) and N-formyl-met-leu-phe (fMLP), but less ROS in response to lipopolysaccharide[171]. Overexpression of constitutively active Rac1 and Cdc42 both significantly increased superoxide anion production in cardiomyocytes[172]. Furthermore, in a human promyelocytic HL-60 cell line, expression of dominant-inhibitory forms of Cdc42 (Cdc42N17) interfered with the NADPH oxidase activation through the GTPloading of Rac and Ras, intracellular calcium mobilization, activation of the p38 MAPK pathway and superoxide production[173]. In platelets, a dual requirement exists for Rac1 and Cdc42 for proper platelet production and function [174,175], yet the manner in which these Rho GTPase family members work together to regulate platelet physiology remains unknown. For instance, while Rac1 and Cdc42 may coordinate platelet formation through spatially and temporally distinct signalling events, or colocalize and work simultaneously to regulate platelet formation, whether Cdc42 and Rac1 coordinate to regulate the NADPH oxidase system and ROS formation remains unknown.

Pleiotropic effects of statins: implications for Rho GTPase-mediated oxidative stress

Platelets play a deleterious role in atherogenesis as well as atherothrombosis, myocardial infarction and ischaemic stroke. Hypercholesterolemia, one of the main driving forces of atherogenesis, also enhances platelet reactivity, which, in turn, promotes vascular inflammation and thrombosis. Oxidized low-density-lipoproteins (oxLDLs) promote platelet hyperactivity and procoagulant phenotype via direct binding to CD36 and LOX1 on the platelet membrane[176]. Pharmacological inhibition or genetic ablation of NOX2 significantly reduced oxLDL-induced ROS formation via tyrosine kinase and PKC signalling in human and murine platelets, respectively [60]. The current standard of care for prevention of cardiovascular events includes pharmacotherapy with 3-hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase inhibitors, more commonly known as statins. Although initially described as lipidlowering drugs, pleiotropic effects of statins were first hypothesized in 2008 by Wang and colleagues [177,178].

To date, the biological mechanism underlying pleiotropic effects of statins, including immunomodulation, anti-inflammation and antioxidation, endothelial protection and pro-angiogenesis, is attributed to inhibition of isoprenoid intermediates [179-181]. Prenylation of proteins is performed by geranylgeranyltransferase type I (GGTase-I) by transferring a 20-carbon geranylgeranyl lipid to a cysteine residue of proteins, including Rac1, RhoA and Cdc42[182]. Such process enhances hydrophobicity and facilitates membrane anchoring of Rho proteins for subcellular targeting, effector binding, GTP loading and activation [182-185]. Statins lower cholesterol levels by blocking mevalonate synthesis, but also inhibit Rho-GTPase isoprenylation through reducing geranyl-geranylpyrophosphate (GGPP), the lipid substrate of GGTase-I, generated during cholesterol biosynthesis and therefore affect downstream Rho effectors such as Rho kinase and NADPH oxidase [186-190].

Historically, targeting GGTase-I has been proposed as a strategy for treating inflammatory and autoimmune disorders, although the idea that blocking Rho protein geranylgeranylation would inhibit inflammation was recently challenged by some studies in GGTase-I-deficient mice. Knockout of GGTase-I catalytic subunit in macrophages eliminated Rho protein geranygeranylation, yet Rac1, RhoA and Cdc42 still accumulated in their GTP-bound active form. Furthermore, p38 and NF-κB activities remained high in GGTase-I-deficient macrophages [191,192]. In another study, GGTase-I-mediated prenylation of proteins was thought to act as a break on innate immune responses in macrophage by limiting Rac1 effector interactions, and therefore blocking prenylation unleashes proinflammatory signalling, even though prenylation is not required for GTP loading and activation of Rho proteins[185]. Furthermore, statin use is associated with elevated coronary calcification in highrisk patients and disrupts the Rac1-Rho GDI complex leading to increased active Rac1 in primary monocytes and macrophages[193].

Nevertheless, statins are more often associated with anti-inflammatory rather than pro-inflammatory effects [194-200], which could be due to the drug action on platelets in addition to lymphocytes and macrophages [201–206]. Influence of statins on platelet function has been investigated in patients with hypercholesterolemia, diabetes mellitus, metabolic syndrome and established atherosclerosis [207-212]. Statins decrease oxidative stress and platelet activation via NOX2 and phospholipase A2 activation, along with inhibition of platelet recruitment, isoprostanes and thromboxane A2 generation [213-215]. Of note, these effects are observed immediately after administration of statin and therefore seem to be independent of lipid lowering[216]. Upon prolonged statin treatment, continued suppression of platelet thromboxane formation has been observed in parallel with lipid lowering[217]. Statin-treated platelets have decreased PAR1 expression, reduced ADP/ATP release at concentrations 10-100-fold lower than therapeutic plasma levels in hypercholesterolemic patients, and Rho GTPase dissociation from Rho GDI and association with platelet membrane, stimulated by thrombin or ADP, has been shown to be inhibited in the presence of statin [210,218]. Statin-treated platelets also failed to enhance oxygen radical production of neutrophils, potentially as a mechanism to protect the vasculature from excessive superoxide anion burden[219]. Furthermore, it is well established that ROS negatively influences NO biosynthesis and activity; therefore, impaired NOX function mediated by statins might up-regulate NO generation, accounting for the antioxidant effects by statins[215]. Statins can also directly enhance platelet cGMP and up-regulate platelet eNOS activity [212,220,221]. Overall, antiplatelet mechanisms by which the NADPH oxidase system is attenuated and subsequent ROS formation is reduced by statins might very well be contributing to statins' efficacy and overall cardiovascular benefit[222].

Conclusions and future directions

Rho GTPases are molecular switches that control numerous signal transduction pathways, from regulating cytoskeletal dynamics, motility and membrane transport, to cellular oxidation processes that altogether play critical roles in platelet function and thrombus formation. Furthermore, emerging cell biological studies continue to find increasing regulatory and functional overlap between the NADPH oxidase system and Rho GTPases in various cell models, including platelets. However, the exact mechanisms by which Rho GTPases orchestrate ROS formation and oxidative stress in a spatio-temporal manner remain to be explored further.

Rho GTPases have long been of interest as therapeutic targets relevant to thrombosis, vascular inflammation and other platelet-associated diseases. Although current literature suggests that the development of therapies targeting platelet Rho GTPases may be beneficial in the context of cardiovascular disease, the

potential off-target effects of systemic inhibition of Rho GTPases have yet to be determined. Whereas Rac1, RhoA and Cdc42 share overlapping signalling pathways, each of them also have specific upstream and downstream regulators that may provide means to identify safe targets for specific disease contexts. For instance, it has recently been shown that the Rhodownstream mediator ROCK2 is a key regulator of platelet activation in thrombosis, but does not mediate the contribution of platelets to atherosclerotic plaque formation and vascular remodelling[223].

As agents targeting specific Rho GTPase interactions with Rho GEFs and other partners emerge further as potential therapeutics, future efforts may fine tune Rho GTPase activities in platelets in specific disease contexts - including ROS generation, where agents against NADPH also show potential[224]. Altogether, basic cell biological studies of signalling in cytoskeletal regulation and oxidative stress in parallel with translational studies of platelet behaviours in the context of statins and other cardiovascular therapeutics are evolving together to improve basic understanding of cell function while driving the development of strategies to target diseases.

Nonstandard Abbreviations and Acronyms

AA	arachidonic acid
ARHGEF1	Rho Guanine Nucleotide Exchange Factor 1
Bcl-2	B-cell lymphoma-2
Bcl-xl	B-cell lymphoma-extra large
CD	cluster of differentiation
Cdc42	cell division control protein 42 homolog
DNA	deoxyribonucleic acid
ER	endoplasmic reticulum
ERK	extracellular signal-regulated kinase
FAD	flavin adenine dinucleotide
FAK	focal adhesion kinase
fMLP	N-Formylmethionyl-leucyl-phenylalanine
GAP	GTPase-activating protein
GEF	guanine nucleotide exchange factor
GDI	guanine dissociate inhibitor
GDP	guanine diphosphate
GGPP	geranyl-geranylpyrophosphate
GGTase-I	geranylgeranyltransferase type I
GP	glycoprotein
GPCR	G-protein coupled receptors
GTP	guanine triphosphate
IMAC	inner membrane anion channel
IP_3	inositol trisphosphate
ITAM	immunoreceptor tyrosine-based activation
	motif
LOX1	lectin-like oxidized low-density lipoprotein
	(LDL) receptor-1
T 70.0	1, 1 1, 1

lipopolysaccharide

mitogen-activated protein kinase

LPS

MAPK

MK2

mitogen-activated protein kinase activated protein kinase 2 (MAPKAPK2) myosin light chain

MYPT myosin phosphatase targeting protein mitochonrial permeability transition pore mPTP **NADH** nicotinamide adenine dinucleotide

nicotinamide adenine dinucleotide phosphate **NADPH**

NF_KB nuclear factor kappa B NOX NADPH oxidase

MLC

oxLDL oxidized low-density lipoprotein

PAK p21-activated kinase protease-activated receptor **PAR** PI3K phosphoinositide-3 kinase

PKA protein kinase A **PKB** protein kinase B **PKC** protein kinase C **PKG** protein kinase G

Rac Ras-related C3 botulinum toxin substrate 1

Rho Ras homolog family member

ROCK Rho-associated coiled-coil containing protein

kinases

ROS reactive oxygen species

RTN4 reticulon 4

Syk spleen associated tyrosine kinase TRAF-4 TNF receptor-associated factor 4

tumor necrosis factor **TNF**

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Disclosure statement

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